# STIC Biotechnology Systems Branch

# RAW SEQUENCE LISTING ERROR REPORT

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:**

Application Serial Number: 10/573,718
Source: IFW P
Date Processed by STIC: 4/6/06

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**
**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221**

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

**http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):** U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

## Raw Sequence Listing Error Summary

| ERROR DETECTED | SUGGESTED CORRECTION | SERIAL NUMBER: 10/573,718 |
|---|---|---|

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

| | | |
|---|---|---|
| 1 | Wrapped Nucleics<br>Wrapped Aminos | The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping." |
| 2 | Invalid Line Length | The rules require that a line **not exceed** 72 characters in length. This includes white spaces. |
| 3 | Misaligned Amino Numbering | The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead. |
| 4 | Non-ASCII | The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.** |
| 5 | Variable Length | Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing. |
| 6 | PatentIn 2.0 "bug" | A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.** |
| 7 | Skipped Sequences (OLD RULES) | Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:<br>(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)<br>(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)<br>(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)<br>This sequence is intentionally skipped<br>Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences. |
| 8 | Skipped Sequences (NEW RULES) | Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.<br><210> sequence id number<br><400> sequence id number<br>000 |
| 9 | Use of n's or Xaa's (NEW RULES) | Use of n's and/or Xaa's have been detected in the Sequence Listing.<br>Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.<br>In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents. |
| 10 | Invalid <213> Response | Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below) |
| 11 | Use of <220> | Sequence(s) ___ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules |
| 12 | PatentIn 2.0 "bug" | Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk. |
| 13 | Misuse of n/Xaa | "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid |

IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/573,718

DATE: 04/06/2006
TIME: 11:00:12

Input Set : A:\08917-116US1 Seq_List.txt
Output Set: N:\CRF4\04062006\J573718.raw

pp 1, 4

Does Not Comply
Corrected Diskette Needed

<110> APPLICANT: Toraya, Tetsuo
Tobimatsu, Takamasa
Yamanishi, Mamoru
Mori, Kouichi
Kajiura, Hideki
Yamada, Seiki
Yuzuki, Michio
Azuma, Muneaki
Hara, Tetsuya
Yasuda, Shinzo
<120> TITLE OF INVENTION: Method for Production of 3-hydroxypropionaldehyde
<130> FILE REFERENCE: 08917-116US1
C--> <140> CURRENT APPLICATION NUMBER: US/10/573,718
C--> <141> CURRENT FILING DATE: 2006-03-27
<150> PRIOR APPLICATION NUMBER: PCT/JP2004/014213
<151> PRIOR FILING DATE: 2004-09-29
<150> PRIOR APPLICATION NUMBER: JP 2003-337663
<151> PRIOR FILING DATE: 2003-09-29
<160> NUMBER OF SEQ ID NOS: 4
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<213> ORGANISM: plasmid pBR322

invalid <213> response. See item 10 on Error Summary Sheet

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PATENT APPLICATION: US/10/573,718 TIME: 11:00:12

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PATENT APPLICATION: US/10/573,718 TIME: 11:00:12

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RAW SEQUENCE LISTING DATE: 04/06/2006
PATENT APPLICATION: US/10/573,718 TIME: 11:00:12

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344 gttcttctcc acgccgctgc cgcccgccgt gttcgcccgc gtctgcgtgg tgaaagcgga   1920
346 cgaactggtg ccgctgcccg gcgatttagc gctggaaaaa gtgcgcgcca ttcgccgcag   1980
348 cgccaaagag cgggtctttg tcaccaacgc cctgcgcgcg ctgcgtcagg tcagccccac   2040
350 cggcaacatt cgcgatattc cgttcgtggt gctggtcggc ggttcgtcgc tggatttcga   2100
352 agtcccgcag ctggtcaccg atgcgctggc gcactaccgc ctggttgccg gacggggaaa   2160
354 tattcgcggc agcgagggcc cccgaaacgc ggtggccacc ggcctgattc tctcctggca   2220
356 taaggagttt gcgcatgaac ggtaatcaca gcgccccggc catcgcgatc gccgtcatcg   2280
358 acggctgcga cggcctgtgg cgcgaagtgc tgctgggtat cgaagaggaa ggtatccctt   2340
360 tccggctcca gcatcacccg gccggagagg tcgtggacag cgcctggcag gcggcgcgca   2400
```

same enzyme as in sequences 3-4, too

same enzyme

RAW SEQUENCE LISTING DATE: 04/06/2006
PATENT APPLICATION: US/10/573,718 TIME: 11:00:12

Input Set : A:\08917-116US1 Seq_List.txt
Output Set: N:\CRF4\04062006\J573718.raw

362 gctcgccgct gctggtgggc atcgcctgcg accgccatat gctggtcgtg cactacaaga 2460
364 atttacccgc atcggcgccg ctttttacgc tgatgcatca tcaggacagt caggcccatc 2520
366 gcaacaccgg taataacgcg gcacggctgg tcaaggggat ccctttccgg gatctgaata 2580
368 gcgaagcaac aggagaacag caggatgaat aagatctcgg gtagcccgcc taatgagcgg 2640
370 gctttttttt atgagaatta caacttatat cgtatggggc tgacttcagg tgctacattt 2700
372 gaagagataa attgcactga aatctagaaa tattttatct gattaataag atgatcttct 2760
374 tgagatcgtt ttggtctgcg cgtaatctct tgctctgaaa acgaaaaaac cgccttgcag 2820
376 ggcggttttt cgaaggttct ctgagctacc aactctttga accgaggtaa ctggcttgga 2880
378 ggagcgcagt caccaaaact tgtcctttca gtttagcctt aaccggcgca tgacttcaag 2940
380 actaactcct ctaaatcaat taccagtggc tgctgccagt ggtgcttttg catgtctttc 3000
382 cgggttggac tcaagacgat agttaccgga taaggcgcag cggtcggact gaacgggggg 3060
384 ttcgtgcata cagtccagct tggagcgaac tgcctacccg gaactgagtg tcaggcgtgg 3120
386 aatgagacaa acgcggccat aacagcggaa tgacaccggt aaaccgaaag gcaggaacag 3180
388 gagagcgcac gagggagccg ccaggggaa acgcctggta tctttatagt cctgtcgggt 3240
390 ttcgccacca ctgatttgag cgtcagattt cgtgatgctt gtcaggggg cggagcctat 3300
392 ggaaaaacgg ctttgccgcg gccctctcac ttccctgtta agtatcttcc tggcatcttc 3360
394 caggaaatct ccgccccgtt cgtaagccat ttccgctcgc cgcagtcgaa cgaccgagcg 3420
396 tagcgagtca gtgagcgagg aagcggaata tatcctgtat cacatattct gctgacgcac 3480
398 cggtgcagcc ttttttctcc tgccacatga agcacttcac tgacaccctc atcagtgcca 3540
400 acatagtaag ccagtataca ctccgctagc gctgatgtcc ggcggtgctt ttgccgttac 3600
402 gcaccacccc gtcagtagct gaacaggagg gacagctgat agaaacagaa gccactggag 3660
404 cacctcaaaa acaccatcat acactaaatc agtaagttgg cagcatcacc cgacgcactt 3720
406 tgcgccgaat aaatacctgt gacggaagat cacttcgcag aataaataaa tcctggtgtc 3780
408 cctgttgata ccgggaagcc ctgggccaac ttttggcgaa aatgagacgt tgatcggcac 3840
410 gtaagaggtt ccaactttca ccataatgaa ataagatcac taccgggcgt atttttgag 3900
412 ttatcgagat tttcaggagc taaggaagct aaaatggaga aaaaaatcac tggatatacc 3960
414 accgttgata tatcccaatg gcatcgtaaa gaacattttg aggcatttca gtcagttgct 4020
416 caatgtacct ataaccagac cgttcagctg gatattacgg ccttttaaa gaccgtaaag 4080
418 aaaaataagc acaagtttta tccggccttt attcacattc ttgcccgcct gatgaatgct 4140
420 catccggaat ttcgtatggc aatgaaagac ggtgagctgg tgatatggga tagtgttcac 4200
422 ccttgttaca ccgttttcca tgagcaaact gaaacgtttt catcgctctg gagtgaatac 4260
424 cacgacgatt tccggcagtt tctacacata tattcgcaag atgtggcgtg ttacggtgaa 4320
426 aacctggcct atttccctaa agggtttatt gagaatatgt ttttcgtctc agccaatccc 4380
428 tgggtgagtt tcaccagttt tgatttaaac gtggccaata tggacaactt cttcgccccc 4440
430 gttttcacca tgggcaaata ttatacgcaa ggcgacaagg tgctgatgcc gctggcgatt 4500
432 caggttcatc atgccgtctg tgatggcttc catgtcggca gaatgcttaa tgaattacaa 4560
434 cagtactgcg atgagtggca gggcggggcg taatttttt aaggcagtta ttggtgccct 4620
436 taaacgcctg gtgctacgcc tgaataagtg ataataagcg gatgaatggc agaaattcga 4680
438 aagcaaattc gacccggtcg tcggttcagg gcagggtcgt taaatagccg cttatgtcta 4740
440 ttgctggttt accggtttat tgactaccgg aagcagtgtg accgtgtgct tctcaaatgc 4800
442 ctgaggccag tttgctcagg ctctccccgt ggaggtaata attgacgata tgatcattta 4860
444 ttctgcctcc caaagcaatt ccgacaccat cgaatggtgc aaaacctttc gcggtatggc 4920
446 atgatagcgc ccggaagaga gtcaattcag ggtggtgaat gtgaaaccag taacgttata 4980
448 cgatgtcgca gagtatgccg gtgtctctta tcagaccgtt tcccgcgtgg tgaaccaggc 5040
450 cagccacgtt tctgcgaaaa cgcgggaaaa agtggaagcg gcgatggcgg agctgaatta 5100
452 cattcccaac cgcgtggcac aacaactggc gggcaaacag tcgttgctga ttggcgttgc 5160
454 cacctccagt ctggccctgc acgcgccgtc gcaaattgtc gcggcgatta aatctcgcgc 5220
456 cgatcaactg ggtgccagcg tggtggtgtc gatggtagaa cgaagcggcg tcgaagcctg 5280
458 taaagcggcg gtgcacaatc ttctcgcgca acgcgtcagt gggctgatca ttaactatcc 5340

**VERIFICATION SUMMARY** DATE: 04/06/2006
PATENT APPLICATION: **US/10/573,718** TIME: 11:00:13

Input Set : **A:\08917-116US1 Seq_List.txt**
Output Set: **N:\CRF4\04062006\J573718.raw**

L:18 M:270 C: Current Application Number differs, Replaced Current Application No
L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date